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<110> Sharp, David J.
Rogers, Gregory C.
Scholey, Jonathon M.

<120> PEPTIDE INHIBITORS OF CELLULAR PROLIFERATION

<130> UC069.001A

<140> 09/782,816
<141> 2001-02-14

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<220>

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Thr Pro Gln Gln Lys Tyr Gln Arg Leu Leu His Glu Val Gln Glu Leu
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<211> 14
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Arg Leu Leu His Glu Val Gln Glu.Leu Thr Thr
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Val Gln Glu Leu Thr Thr
1 5

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<210> 23
<211> 6
<212> PRT
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<220>

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1 5

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<212> PRT
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<220>

<223> Peptide sequence

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<212> PRT
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1 5 10 15
Met Asn Glu Leu Leu Asn
20

<210> 27
<211> 21
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<213> Unknown

<220>

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Glu Lys Glu Thr Pro Val Gln Lys Cys Gln Arg Leu Gln Ile Glu Met
1 5 10 15
Asn Glu Leu Leu Asn
20

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<213> Unknown

<220>

<223> Peptide sequence

<400> 28
Lys Glu Thr Pro Val Gln Lys Cys Gln Arg Leu Gln Ile Glu Met Asn
1 5 10 15
Glu Leu Leu Asn
20

<210> 29
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<400> 29
Glu Thr Pro Val Gln Lys Cys Gln Arg Leu Gln Ile Glu Met Asn Glu
1 5 10 15
Leu Leu Asn

<210> 30
<211> 18
<212> PRT
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<220>

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<400> 30
Thr Pro Val Gln Lys Cys Gln Arg Leu Gln Ile Glu Met Asn Glu Leu
1 5 10 15
Leu Asn

<210> 31
<211> 17
<212> PRT

<213> Unknown

<220>

<223> Peptide sequence

<400> 31
Pro Val Gln Lys Cys Gln Arg Leu Gln Ile Glu Met Asn Glu Leu Leu
 1               5                  10                  15
Asn

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Val Gln Lys Cys Gln Arg Leu Gln Ile Glu Met Asn Glu Leu Leu Asn
 1               5                  10                  15

<210> 33
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<212> PRT
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<220>

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Gln Lys Cys Gln Arg Leu Gln Ile Glu Met Asn Glu Leu Leu Asn
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<210> 34
<211> 14
<212> PRT
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<220>

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Lys Cys Gln Arg Leu Gln Ile Glu Met Asn Glu Leu Leu Asn
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Cys Gln Arg Leu Gln Ile Glu Met Asn Glu Leu Leu Asn
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<212> PRT
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<400> 36
Gln Arg Leu Gln Ile Glu Met Asn Glu Leu Leu Asn
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<220>

<223> Peptide sequence

<400> 37
Arg Leu Gln Ile Glu Met Asn Glu Leu Leu Asn
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<210> 38
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<400> 38
Leu Gln Ile Glu Met Asn Glu Leu Leu Asn
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Gln Ile Glu Met Asn Glu Leu Leu Asn
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<400> 41
Glu Met Asn Glu Leu Leu Asn
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<400> 43
Asn Glu Leu Leu Asn
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<210> 44
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Val Ala Thr Val Ile Ser Thr Ala Arg
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<210> 46
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Val Ala Thr Val Ile Ser Thr Ala
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Val Ala Thr Val Ile Ser Thr
 1               5

<210> 48
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Val Ala Thr Val Ile Ser
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Val Ala Thr Val Ile
 1               5

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<223> Peptide sequence

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            20                  25                  30
Glu Ser Ala Thr Glu Glu Lys Leu Thr Pro Val Xaa Leu Ala Lys Gln
        35                  40                  45
Leu Ala Ala Leu
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<223> Peptide sequence

<400> 52
Gly Glu Lys Glu Thr Pro Val Gln Lys Cys Gln Arg Leu Gln Ile Glu
 1               5                  10                  15
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            20                  25                  30
Val Ala Asp Glu Glu Lys Gln Ser Tyr Asp Ala Val Val Ala Thr Val
        35                  40                  45
Ile Ser Thr Ala Arg
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<210> 53
<211> 406
<212> PRT
<213> Homo sapiens

<400> 53
Met Ala Asp Pro Lys Tyr Ala Asp Leu Pro Gly Ile Ala Arg Asn Glu
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            20                  25                  30
Phe Asp Ala Phe Ala Gln Glu Leu Glu Glu Leu Thr Ser Thr Ser Val
        35                  40                  45
Glu His Ile Ile Val Asn Pro Asn Ala Ala Tyr Asp Lys Phe Lys Asp
    50                  55                  60
Lys Arg Val Gly Thr Lys Gly Leu Asp Phe Ser Asp Arg Ile Gly Lys
65                  70                  75                  80
Thr Lys Arg Thr Gly Tyr Glu Ser Gly Glu Tyr Glu Met Leu Gly Glu
                85                  90                  95
Gly Leu Gly Val Lys Glu Thr Pro Gln Gln Lys Tyr Gln Arg Leu Leu
            100                 105                 110

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His Glu Val Gln Glu Leu Thr Thr Glu Val Glu Lys Ile Lys Thr Thr
        115                 120                 125
Val Lys Glu Ser Ala Thr Glu Glu Lys Leu Thr Pro Val Leu Leu Ala
    130                 135                 140
Lys Gln Leu Ala Ala Leu Lys Gln Gln Leu Val Ala Ser His Leu Glu
145                 150                 155                 160
Lys Leu Leu Gly Pro Asp Ala Ala Ile Asn Leu Thr Asp Pro Asp Gly
                165                 170                 175
Ala Leu Ala Lys Arg Leu Leu Leu Gln Leu Glu Ala Thr Lys Asn Ser
            180                 185                 190
Lys Gly Gly Ser Gly Gly Lys Thr Thr Gly Thr Pro Pro Asp Ser Ser
        195                 200                 205
Leu Val Thr Tyr Glu Leu His Ser Arg Pro Glu Gln Asp Lys Phe Ser
    210                 215                 220
Gln Ala Ala Lys Val Ala Glu Leu Glu Lys Arg Leu Thr Glu Leu Glu
225                 230                 235                 240
Thr Ala Val Arg Cys Asp Gln Asp Ala Gln Asn Pro Leu Ser Ala Gly
                245                 250                 255
Leu Gln Gly Ala Cys Leu Met Glu Thr Val Glu Leu Leu Gln Ala Lys
            260                 265                 270
Val Ser Ala Leu Asp Leu Ala Val Leu Asp Gln Val Glu Ala Arg Leu
        275                 280                 285
Gln Ser Val Leu Gly Lys Val Asn Glu Ile Ala Lys His Lys Ala Ser
    290                 295                 300
Val Glu Asp Ala Asp Thr Gln Ser Lys Val His Gln Leu Tyr Glu Thr
305                 310                 315                 320
Ile Gln Arg Trp Ser Pro Ile Ala Ser Thr Leu Pro Glu Leu Val Gln
                325                 330                 335
Arg Leu Val Thr Ile Lys Gln Leu His Glu Gln Ala Met Gln Phe Gly
            340                 345                 350
Gln Leu Leu Thr His Leu Asp Thr Thr Gln Gln Met Ile Ala Asn Ser
        355                 360                 365
Leu Lys Asp Asn Thr Thr Leu Leu Thr Gln Val Gln Thr Thr Met Arg
    370                 375                 380
Glu Asn Leu Ala Thr Val Glu Gly Asn Phe Ala Ser Ile Asp Glu Arg
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Met Lys Lys Leu Gly Lys
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<213> Mouse

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            20                  25                  30
Phe Asp Ala Glu Glu Leu Ser Ser Thr Ser Val Glu His Ile Ile Val
        35                  40                  45
Asn Pro Asn Ala Ala Tyr Asp Lys Phe Lys Asp Lys Arg Val Gly Thr
    50                  55                  60
Lys Gly Leu Asp Phe Ser Asp Arg Ile Gly Lys Thr Lys Arg Thr Gly
65                  70                  75                  80
Tyr Glu Ser Gly Asp Tyr Glu Met Leu Gly Glu Gly Leu Gly Val Lys
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                85                  90                  95
Glu Thr Pro Gln Gln Lys Tyr Gln Arg Leu Leu His Glu Val Gln Glu
            100                 105                 110
Leu Thr Thr Glu Val Glu Lys Ile Lys Thr Thr Val Lys Glu Ser Ala
        115                 120                 125
Thr Glu Glu Lys Leu Thr Pro Val Val Leu Ala Lys Gln Leu Ala Ala
    130                 135                 140
Leu Lys Gln Gln Leu Val Ala Ser His Leu Glu Lys Leu Leu Gly Pro
145                 150                 155                 160
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<213> Drosophila melanogaster

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Tyr Tyr Glu Glu Glu Pro Glu Asn Glu Ala Ile Glu Arg Leu His Ile
        35                  40                  45
Ser Pro Ser Val Ala His Lys Arg Phe Ser Gly Ala Thr Val Glu Gly
    50                  55                  60
Ser Val Asp Phe Thr Asp Arg Ile Gly Arg Arg Met Cys Arg Gly Tyr
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65                  70                  75                  80
Asp Thr Arg Gly Ser Ser Asp Tyr Glu Leu Val Gly Gln Gly Glu Lys
                85                  90                  95
Glu Thr Pro Val Gln Lys Cys Gln Arg Leu Gln Ile Glu Met Asn Glu
            100                 105                 110
Leu Leu Asn Glu Val Ala Ala Leu Gln Val Asp Arg Lys Val Ala Asp
        115                 120                 125
Glu Glu Lys Gln Ser Tyr Asp Ala Val Ala Thr Val Ile Ser Thr Ala
    130                 135                 140
Arg Lys Val Leu Glu Ser Leu Lys Leu Glu Gln Val Leu Gly Lys Glu
145                 150                 155                 160
Gln Thr Pro Gly Ser Lys Gln Val Lys Ala Leu Ile Ser Gln Val Glu
                165                 170                 175
Glu Phe Lys Gln Ser Gly Val Leu Thr Ala Ile Pro Thr Pro Gly Thr
            180                 185                 190
Asp Leu Ala Ala Thr Ala Arg Val Ala Ser Leu Glu Gln Arg Ile Ser
        195                 200                 205
Gln Leu Glu Lys Val Leu Gly Ala Gln Pro Asp Lys Leu Ser Arg Leu
    210                 215                 220
Thr Ala Ala Thr Asn Thr Thr Asn Val Leu Glu Ala Val Arg His Leu
225                 230                 235                 240
Ser Thr Lys Ala Ala Leu Ile Gln Pro Asp Lys Leu Asp Thr Ile Glu
                245                 250                 255
Gln Arg Leu Thr Ser Leu Ala Gly Lys Met Asp Ala Ile Ala Glu Lys
            260                 265                 270
Ser Ser Gly Ser Ala Gln Asp Ala Lys Arg Asp Gln Lys Ile Thr Glu
        275                 280                 285
Leu Tyr Asp Ile Ala Lys Arg Thr Glu Pro Val Val Glu Ile Leu Pro
    290                 295                 300
His Val Ile Glu Arg Met Gln Ala Leu Glu Ala Leu His Lys Tyr Ala
305                 310                 315                 320
Asn Asn Phe Ala Lys Ile Ile Ala Glu Ile Glu Gln Lys Gln Gly Thr
                325                 330                 335
Ile Thr Thr Ser Leu Val Asn Asn Lys Glu Leu Leu His Ser Val Gln
            340                 345                 350
Glu Thr Phe Ala Gln Asn Leu Glu Thr Ile Asn Ser Lys Val Ala Lys
        355                 360                 365
Val Glu Gln Arg Val Ala Ala Ile Ser Ser Ala Lys
    370                 375                 380
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